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# NOTICE OF ALLOWANCE AND FEE(S) DUE

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07/23/2008

THE MCCALLUM LAW FIRM, P. C. 685 BRIGGS STREET PO BOX 929 ERIE. CO 80516

EXAMINER

MOORE, WILLIAM W

ART UNIT PAPER NUMBER

1656

DATE MAILED: 07/23/2008

APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/554,237	08/04/2006	Kerry Michelle Dunse	007193-13	1348

TITLE OF INVENTION: INSECT CHYMOTRYPSIN AND INHIBITORS THEREOF

APPLN. TYPE	SMALL ENTITY	ISSUE FEE DUE	PUBLICATION FEE DUE	PREV. PAID ISSUE FEE	TOTAL FEE(S) DUE	DATE DUE
nonprovisional	NO	\$1440	\$300	\$0	\$1740	10/23/2008

THE APPLICATION IDENTIFIED ABOVE HAS BEEN EXAMINED AND IS ALLOWED FOR ISSUANCE AS A PATENT. PROSECUTION ON THE MERITS IS CLOSED. THIS NOTICE OF ALLOWANCE IS NOT A GRANT OF PATENT RIGHTS. THIS APPLICATION IS SUBJECT TO WITHDRAWAL FROM ISSUE AT THE INITIATIVE OF THE OFFICE OR UPON PETITION BY THE APPLICANT. SEE 37 CFR 1.313 AND MPEP 1308.

THE ISSUE FEE AND PUBLICATION FEE (IF REQUIRED) MUST BE PAID WITHIN THREE MONTHS FROM THE MAILING DATE OF THIS NOTICE OR THIS APPLICATION SHALL BE REGARDED AS ABANDONED. THIS STATUTORY PERIOD CANNOT BE EXTENDED. SEE 35 U.S.C. 151. THE ISSUE FEE DUE INDICATED ABOVE DOES NOT REFLECT A CREDIT FOR ANY PREVIOUSLY PAID ISSUE FEE IN THIS APPLICATION. IF AN ISSUE FEE HAS PREVIOUSLY BEEN PAID IN THIS APPLICATION (AS SHOWN ABOVE), THE RETURN OF PART B OF THIS FORM WILL BE CONSIDERED A REQUEST TO REAPPLY THE PREVIOUSLY PAID ISSUE FEE TOWARD THE ISSUE FEE NOW DUE.

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								(Signature)
								(Date)
APPLICATION NO.	FILING DATE		FIRST NAMED INVEN	TOR		ATTOI	RNEY DOCKET NO.	CONFIRMATION NO.
10/554,237 ITLE OF INVENTION	08/04/2006 :: INSECT CHYMOTRY	PSIN AND INHIBITOR	Kerry Michelle Dun S THEREOF	ise			007193-13	1348
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nonprovisional	NO	\$1440	\$300		\$0		\$1740	10/23/2008
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MOORE, W	ILLIAM W	1656	530-370000					
Change of corresponde FR 1.363).  Change of corresp Address form PTO/SI  "Fee Address" ind PTO/SB/47; Rev 03-0 Number is required.	(1) the names of u or agents OR, alter (2) the name of a s registered attorney 2 registered patent	printing on the patent front page, list names of up to 3 registered patent attorneys ats OR, alternatively, name of a single firm (having as a member a red attorney or agent) and the names of up to tered patent attorneys or agents. If no name is no name will be printed.						
PLEASE NOTE: Unl recordation as set fort (A) NAME OF ASSIG	less an assignee is ident h in 37 CFR 3.11. Comp GNEE		data will appear on the Tasubstitute for filing (B) RESIDENCE: (C)	he pa g an a	tent. If an assigned ssignment. and STATE OR CO	OUNT	RY)	cument has been filed for
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36234 75	590 07/23/2008		EXAM	INER	
THE MCCALLU	JM LAW FIRM, P. C	MOORE, WILLIAM W			
685 BRIGGS STR	EET		ART UNIT	PAPER NUMBER	
PO BOX 929 ERIE, CO 80516			1656 DATE MAILED: 07/23/200	8	

## Determination of Patent Term Adjustment under 35 U.S.C. 154 (b)

(application filed on or after May 29, 2000)

The Patent Term Adjustment to date is 112 day(s). If the issue fee is paid on the date that is three months after the mailing date of this notice and the patent issues on the Tuesday before the date that is 28 weeks (six and a half months) after the mailing date of this notice, the Patent Term Adjustment will be 112 day(s).

If a Continued Prosecution Application (CPA) was filed in the above-identified application, the filing date that determines Patent Term Adjustment is the filing date of the most recent CPA.

Applicant will be able to obtain more detailed information by accessing the Patent Application Information Retrieval (PAIR) WEB site (http://pair.uspto.gov).

Any questions regarding the Patent Term Extension or Adjustment determination should be directed to the Office of Patent Legal Administration at (571)-272-7702. Questions relating to issue and publication fee payments should be directed to the Customer Service Center of the Office of Patent Publication at 1-(888)-786-0101 or (571)-272-4200.

	Application No.	Applicant(s)	
	10/554,237	DUNSE ET AL.	
Notice of Allowability	Examiner	Art Unit	
	   WILLIAM W. MOORE	1656	
The MAILING DATE of this communication apperature All claims being allowable, PROSECUTION ON THE MERITS IS herewith (or previously mailed), a Notice of Allowance (PTOL-85) NOTICE OF ALLOWABILITY IS NOT A GRANT OF PATENT RIOF the Office or upon petition by the applicant. See 37 CFR 1.313	(OR REMAINS) CLOSED in this ap or other appropriate communication IGHTS. This application is subject to	plication. If not included n will be mailed in due course. <b>THIS</b>	
1. $\boxtimes$ This communication is responsive to <u>the election made 18</u>	April 2008 and the interview condu	<u>cted 2 July 2008</u> .	
2. 🔀 The allowed claim(s) is/are <u>68-86</u> .			
<ul> <li>3.</li></ul>	e been received. e been received in Application No		
<ol><li>Copies of the certified copies of the priority do</li></ol>	cuments have been received in this	national stage application from the	
International Bureau (PCT Rule 17.2(a)).  * Certified copies not received:			
Applicant has THREE MONTHS FROM THE "MAILING DATE" noted below. Failure to timely comply will result in ABANDONN THIS THREE-MONTH PERIOD IS NOT EXTENDABLE.  4. A SUBSTITUTE OATH OR DECLARATION must be subm INFORMAL PATENT APPLICATION (PTO-152) which give	MENT of this application.  itted. Note the attached EXAMINER	R'S AMENDMENT or NOTICE OF	
5. CORRECTED DRAWINGS ( as "replacement sheets") mus	st be submitted.		
(a) ☐ including changes required by the Notice of Draftspers		-948) attached	
1)  hereto or 2)  to Paper No./Mail Date	•	,	
(b) ☐ including changes required by the attached Examiner's Paper No./Mail Date  Identifying indicia such as the application number (see 37 CFR 1 each sheet. Replacement sheet(s) should be labeled as such in t	.84(c)) should be written on the drawi	ings in the front (not the back) of	
DEPOSIT OF and/or INFORMATION about the depo attached Examiner's comment regarding REQUIREMENT	sit of BIOLOGICAL MATERIAL	must be submitted. Note the	
Attachment(s)			
1. Notice of References Cited (PTO-892)	5. Notice of Informal F	Patent Application	
2. Notice of Draftperson's Patent Drawing Review (PTO-948)	6.  ☐ Interview Summary Paper No./Mail Da		
3. Information Disclosure Statements (PTO/SB/08), Paper No./Mail Date 20070316 & 20070412	7. 🛛 Examiner's Amend		
Examiner's Comment Regarding Requirement for Deposit of Biological Material	<ul><li>8. ☑ Examiner's Statem</li><li>9. ☐ Other</li></ul>	ent of Reasons for Allowance	

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#### **EXAMINER'S AMENDMENT**

An examiner's amendment to the record appears below. Should the changes and/or additions be unacceptable to applicant, an amendment may be filed as provided by 37 CFR 1.312. To ensure consideration of such an amendment, it MUST be submitted no later than the payment of the issue fee.

## Amend the two paragraphs at page 84, lines 2-19, of the specification thus:

The cDNA clones were grouped on the basis of restriction fragment patterns obtained using combinations of the endonucleases <code>BamHI</code>, <code>XhoII</code>, <code>KpnI</code>, <code>SacII</code>, <code>SacII</code>, <code>and SalI</code> (Promega). RT-PCR products and cDNA inserts were sequenced in both directions using M13 universal primers at either Micromon sequencing facility at Monash University (Melbourne) or SUPAMAC at the Royal Prince Alfred Hospital in Sydney. The sequence data was edited using the BioEdit v5.0.9.1 software written by Tom Hall, North Carolina State University freely available at the web address: <a href="mbio.ncsu.edu/BioEdit/bioedit.html">mbio.ncsu.edu/BioEdit/bioedit.html</a>. Sequence homologies were assessed using the BLASTN search facility at National Centre for Biotechnology Information (NCBI) and further multiple sequence alignments were performed using ClustalW version 1.4. at the Network

BLASTN search facility at National Centre for Biotechnology Information (NCBI) and further multiple sequence alignments were performed using ClustalW version 1.4. at the Network Protein Sequence Analysis facility (NPSA; http://npsa-pbil.ibcp.fr/cgi-bin/align\_clustalw.pl.) (Combet et al., TIBS. 25: 147-150, 2000).

The web based program 'PSORT II' available at the Human Genome Centre at the University of Tokyo (http://psort.nibb.ac.jp/form2.html), was used to predict signal peptide cleavage points. UTRscan was used to detect functional elements in the 3' untranslated regions of the cDNA clones [Pesole, *Trends Genet*, **15**: 378, 1999]. (http://bighost.area.ba.enr.it/BIG/UTRScan/).

## Amend the paragraph at page 85, lines 9-25, of the specification thus:

The deduced amino acid sequences from the cDNA clones HpF2B (sensitive) and HpF5 (insensitive) were modeled on the structures of the *Bos taurus* (bovine) and fire ant chymotrypsins, obtained from the Research Collaboratory for Structural Bioinformatics (RCSB) Protein Data Bank site (http://www.rcsb.org/pdb/). The *Helicoverpa* chymotrypsins are predicted to adopt similar structures to those reported for all the chymotrypsin structures available in the PDB databank. The modeled structure consists of the classic serine protease fold consisting of two, six-stranded anti-parallel β barrels with the catalytic triad located between the two domains. Two surface loops, 60 and 142 are considerably larger in the *H. punctigera* chymotrypsins (Figures 15 and 16). Due to the limitations of modelling, a small amount of ambiguity was present in several surface loops, some of which are cleaved in mammalian chymotrypsins (loop 142), but remain intact within insect chymotrypsins. The only reported crystal structure of an insect ehymotrypsin is from the fire ant, *Soenopsis invicta* (Botos *et al., Journal of Molecular Biology 298:* 895-901, 2000) and this was used to help refine the orientation of the surface loops on the model of the *Helicoverpa* chymotrypsin.

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To assist the printer, the non-amended claims 70 and 71 presented by Applicant in the amendment filed 18 April 2008 are reproduced following the amended claim 69 with the designation "Previously presented".

Cancel claims 27-51 and 55-68.

### Amend claim 69 thus:

- 69. (Amended) An antagonist that comprises an amino acid sequence having at least 90% amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO:81 and that inhibits the proteolytic activity of the approtein having the approacid sequence set forth in selected from the group consisting of SEQ ID NO:2[[,]] wherein said protein exhibits resistance to a protease inhibitor (PI) from *Nicotiana alata*, and wherein said antagonist comprises the amino acid sequence set forth in SEQ ID NO:81 or an amino acid sequence having at least 90% amino acid sequence identity thereto.
- 70. (Previously presented) The antagonist of Claim 69 wherein said antagonist comprises the amino acid sequence set forth in SEQ ID NO:81.
- 71. (Previously presented) A composition comprising an antagonist of Claim 69.

## Add the new claims 72-83:

- 72. (New) The antagonist of Claim 69 wherein said antagonist comprises an amino acid sequence having at least 95% amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO:81.
- 73. (New) A composition comprising the antagonist of Claim 72.
- 74. (New) A composition comprising the antagonist of Claim 70.
- 75. (New) An isolated nucleic acid molecule having a sequence encoding the antagonist amino acid sequence according to claim 69.
- 76. (New) The isolated nucleic acid molecule of claim 75, wherein the antagonist comprises an amino acid sequence with at least 95% sequence identity to the amino acid sequence set forth in SEQ ID NO:81.
- 77. (New) The isolated nucleic acid molecule of claim 75, wherein the antagonist comprises the amino acid sequence set forth in SEQ ID NO:81.
- 78. (New) A vector comprising the antagonist-encoding nucleic acid sequence according to claim 75.
- 79. (New) A vector comprising the antagonist-encoding nucleic acid sequence according to claim 76.
- 80. (New) A vector comprising the antagonist-encoding nucleic acid sequence according to claim 77.

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81. (New) An isolated genetically modified cell comprising a nucleic acid molecule having a sequence encoding an amino acid sequence having at least 90% amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO:81 and that inhibits the proteolytic activity of the protein having the amino acid sequence set forth in SEQ ID NO:2.

- 82. (New) An isolated genetically modified cell comprising a nucleic acid molecule having a sequence encoding an amino acid sequence having at least 95% amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO:81 and that inhibits the proteolytic activity of the protein having the amino acid sequence set forth in SEQ ID NO:2.
- 83. (New) An isolated genetically modified cell comprising a nucleic acid molecule having a sequence encoding the amino acid sequence set forth in SEQ ID NO:81.
- 84. (New) The isolated cell of claim 81 which is a genetically modified plant cell.
- 85. (New) The isolated cell of claim 82 which is a genetically modified plant cell.
- 86. (New) The isolated cell of claim 83 which is a genetically modified plant cell.

Authorization for this examiner's amendment was given in a telephone interview with Ms. Donna M. Ferber on 7 July 2008.

The following is an examiner's statement of reasons for allowance:

The non-elected claims are canceled in the above amendment and claim 69 is amended to clarify its intended subject matter. The teachings of the specification at page 24, lines 1-5, page 47, lines 14-19, page 48, lines 19-31, page 93, lines 1-15 and Figure 26, pertaining to the StPotIA inhibitor, and the teachings of Beuning et al., 1994, made of record with Applicant's IDS, at pages 647-652, including their alignment of inhibitor amino acid sequences in Table 2 at page 648, demonstrate the state of the art at the time the invention was made. The teachings of Beuning et al. show (i) that one of skill in the art would readily identify the P<sub>10</sub> through P-8 positions of the active site loop within SEQ ID NO:81 wherein the relatively non-conservative substitutions of proline at the P<sub>1</sub> position of the protease-binding loop, where alanine or threonine occur in other potato PI inhibitors, and alanine at the P-1 position of the protease binding loop, where methionine or leucine occur in other potato PI inhibitors, are likely to contribute to the particular affinity of the inhibitor for the protease of SEQ ID NO:2 and (ii) that such an artisan would be well aware of the need for conserving the two cysteines, as well as the PI family I "signature" structural motif, 2 in modifying SEQ NO:81, where fourteen other positions, including the P9 position are shown by Beuning et al. to be variable in other potato PI

Positions 78-95 of SEQ ID NO:81.

<sup>&</sup>lt;sup>2</sup> Positions 51-63 of SEQ ID NO:81.

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inhibitors' carboxyl-proximal regions and Beuning et al. teach that the amino-proximal region of these protease inhibitors may be truncated in part without affecting protease inhibitory activity. Where the specification both discloses four structurally similar species of inhibitors – in SEQ IDs NOs: 77, 78, 80, and 82 – and the prior art discloses still other similar inhibitors sharing as much as 79% identity with SEQ ID NO:81, and discloses an assay that particularly identifies inhibitors of the proteolytic activity of SEQ ID NO:2, the teachings of the specification and the prior art are adequate to support the alteration of as many as eleven amino acid positions in SEQ ID NO:81. The new claims 72-86 describe subject matter supported by the disclosures at, *inter alia*, page 58, lines 20-31, and page 62, lines 22-28, of the specification.

The priority date for the disclosure of SEQ ID NO:81 herein is the 23 April 2003 filing date of the parent application 60/465,054. The prior art made of record herewith does not disclose or suggest the discovery or preparation of a protease inhibitor, i.e., an antagonist, having an amino acid sequence that shares 90% sequence identity with SEQ ID NO:81 herein, nor does it disclose a polynucleotide encoding such an inhibitor, a vector comprising or an encoded amino acid sequence, that has 90% sequence identity with amino acid sequence set forth in SEQ ID NO:81. Thus claims 69-86 are allowed herewith.

Any comments considered necessary by applicant must be submitted no later than the payment of the issue fee and, to avoid processing delays, should preferably accompany the issue fee. Such submissions should be clearly labeled "Comments on Statement of Reasons for Allowance."

#### Conclusion

Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see http://pair-direct.uspto.gov. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

Any inquiry concerning this communication or earlier communications from the examiner should be directed to William W. Moore whose telephone number is 571.272.0933 and whose FAX number is 571.273.0933. The examiner can normally be reached Monday through Friday between 9:00AM and 5:30PM EST. If attempts to reach the examiner by telephone are unsuccessful, the examiner's Supervisory Primary Examiner, Dr. Kathleen Kerr Bragdon, can be reached at 571.272.0931. The official FAX number for all communications for the organization where this application or proceeding is assigned is 571.273.8300. Any inquiry of a general

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Art Unit: 1652

nature or relating to the status of this application or proceeding should be directed to the receptionist whose telephone number is 571.272.1600.

/Nashaat T. Nashed/ Nashaat T. Nashed, Ph.D. Supervisory Primary Examiner Art Unit 1652

/William W. Moore/ William W. Moore 7 July 2008